



FIG. 1

>SEQID#1_SGP037
ATGTCAACAGCTGCCCTTAATTACTTTGGTCAGAAGTGGTGGGAACCAAGGTGAGAAGGAGAGTGTCTAAGCTCCCGCCTGCTGCAGGACGACA
GGCGGGTGACACCCACAGCTGCCACAGCTCCACTTCAGAGCCTAGGTGTTCTCGGTTTGACCCAGATGGTAGTGGGAGTCCAGCTACCTGGGACAA
TTTTGGGATCTGGGATAACCGCATTGATGAGCCAATTCTGCTGCCACCCAGCATTAAAGTATGGCAAGCCAATTCCTCAAAATCAGCTTGGAAAATG
TGGGGTGGCCTCACAGATTGGCAACGGAAGAGAAATGAAGATCGGTTTGACTTCGCTCAGCTGACAGATGAGGTCTCTGTACTTTGCAGTGTAT
TGATGGACACGGTGGACCTGCAGCAGCTGATTTCTGTCTATACCCACATGGAGAAATGTATTATGGATTTGCTTCTAAGGAGAAGAACTTGGAA
ACTCTGTTGACCTTGGCTTTTCTAGAAATAGATAAGCCTTTTCGAGTCTATGCCCGCCTGCTGCTGATGCAACTCTTCTGACCTCTGGGACTACT
GCAACAGTAGCCCTATTGCGAGATGGTATTGAACCTGGTTGTAGCCAGTGTGGGGACAGCCGGGCTATTTGTGTAGAAAAGGAAAACCCATGA
AGCTGACCATTGACCATACTCCAGAAAGAAAAGATGAAAAGAAAGGATCAAGAAATGTGGTGGTTTTGTAGCTTGGAAATAGTTGGGGCAGC
CTCAGGTAATGGCAGGCTTGCAATGACAAGAAGTATTGGAGATTTGGACCTTAAGACCAGTGGTGTCTATGACAGAACCTGAACCTAAGAGGAT
TAAGTTACATCATGCTGATGACAGCTTCTTGGTCTCACCACAGATGGAAATTAACCTCATGGTGAATAGTCAAGAGATTTGTGACTTTTGTCATC
AGTGCCATGATCCCAACGAAGCAGCCCATGCGGTGACTGAACAGGCAATACAGTACGGTACTGAGGATACAGTACTGCAGTAGTAGTGCCCTTT
TGGTGCCTGGGGAAAAATATAAGAACTCTGAATCAACTTCTCATTACAGCAGAAGCTTTGCTCCAGTGGACGATGGGCCTGA

FIG. 2

>SEQID#2_SGP037
MSTAALITLVRSGNQVRRRVLLSSRLLQDDRRVTPTCHSSTSEPRCSRFPDGGSPATWDFGTDNRIDEPIILLPPSIKYGKPIPKISLENVGCASQIG
KRKENEDRFDAQLTDEVLYFAVYDGHGGPAAADFCHTHMEKIMDLLPKEKNILETLLTAFLEIDKAFSSHARLSADATLLTSGTTATVALLRDGIEL
VVASVGDSRAILCRKGKPNKLTIDHTPERKDEKERIKKCGGFVAWNSLGQPHVNGRLAMTRSIGDLDLKTSGVIAEPETKRKIKLHHADDSFLVLTDDGI
NFMVNSQEICDFVNQCHDPNEAAHAVTEQAIOYGTEDNSTAVVVPFGAWGKYKNSINFSSRSFSSGRWA